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GENERAL INFORMATION:
APPLICANT: WOOD, David G.
TITLE OF INVENTION: Amino acid and derivatives of Glycyl, Norleucyl
TITLE OF INVENTION: Polymorphisms In The Human Genome
FILE REFERENCE: 01/00/0125
CURRENT FILING DATE: 2002-02-21
FUTURE FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 095
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO: 26569
LENGTH: 666
ORGANISM: Human
US 09 028 207A 1982

Query Match:
Best Local Similarity: 100.0% Ident. No. 460033
Matches: 18; Conserved: 0; Mismatches: 0; Indels: 0; Gaps: 0

27 4 AAAAAAGAAAGTAAAGG 21
11111111111111111111
18 1144 AAAAGAAAGTAAAGG 11

GENERAL INFORMATION:
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1 OTHER INFORMATION: n equals a, t, c, g, or r
 2 FEATURE:
 3 NAME/KEY: misc feature
 4 LOCATION: (44)
 5 OTHER INFORMATION: n equals a, t, c, g, or r
 6 FEATURE:
 7 NAME/KEY: misc feature
 8 LOCATION: (44)
 9 OTHER INFORMATION: n equals a, t, c, g, or r
 10 SEQ ID NO: 64

Query Match
 Best Local Similarity 84.8% Score 17.89 Len 13 Length 55%
 Matches 19 Conserved 0 Mismatches 2 Indels 0 Gaps 0

27 1 ATGAAAATGAAAATGAAAAT 21
 28 255 ATGAAAATGAAAATGAAAAT 274

FEATURE:
 US ID: 221 274 6876/

1 Sequence 6876 Application 35/1-221-274
 2 GENERAL INFORMATION:
 3 APPLICATION: Biosoft Inc
 4 TITLE OF INVENTION: New 1 Nucleic Acids and Polypeptides
 5 FILE REFERENCE: 21,274,614
 6 CURRENT APPLICATION NUMBER: 02/000,006
 7 PRIOR APPLICATION NUMBER: 02/000,006
 8 PRIOR FILING DATE: 2000-09-19
 9 PRIOR APPLICATION NUMBER: 02/000,006
 10 PRIOR FILING DATE: 2000-09-19
 11 NUMBER OF SEQ ID NOS: 1,246
 12 SOFTWARE: custom
 13 SEQ ID NO: 64
 14 LENGTH: 64
 15 TYPE: DNA
 16 ORGANISM: Homo sapiens
 17 FEATURE:
 18 NAME/KEY: misc feature
 19 LOCATION: (1)-(64)
 20 OTHER INFORMATION: n equals a, t, c, g, or r
 21 SEQ ID NO: 64

Query Match
 Best Local Similarity 84.8% Score 17.89 Len 13 Length 64%
 Matches 19 Conserved 0 Mismatches 2 Indels 0 Gaps 0

27 1 ATGAAAATGAAAATGAAAAT 21
 28 255 ATGAAAATGAAAATGAAAAT 500

FEATURE:
 US ID: 221 500 19776
 1 Sequence 1977 Application 35/1-221-500
 2 GENERAL INFORMATION:
 3 APPLICATION: Biosoft Inc
 4 TITLE OF INVENTION: Nucleic Acids, Proteins, and Antisense
 5 FILE REFERENCE: 21,274,614
 6 CURRENT APPLICATION NUMBER: 02/000,006
 7 PRIOR APPLICATION NUMBER: 02/000,006
 8 PRIOR FILING DATE: 2000-09-19
 9 PRIOR APPLICATION NUMBER: 02/000,006
 10 PRIOR FILING DATE: 2000-09-19
 11 NUMBER OF SEQ ID NOS: 1,246
 12 SOFTWARE: custom
 13 SEQ ID NO: 197
 14 LENGTH: 1001

1 TYPE: DNA
 2 ORGANISM: Homo sapiens
 3 FEATURE:
 4 NAME/KEY: misc feature
 5 LOCATION: (98)
 6 OTHER INFORMATION: n equals a, t, c, g, or r
 7 SEQ ID NO: 94

Query Match
 Best Local Similarity 84.8% Score 17.89 Len 13 Length 94%
 Matches 19 Conserved 0 Mismatches 2 Indels 0 Gaps 0

27 1 ATGAAAATGAAAATGAAAAT 21
 28 505 ATGAAAATGAAAATGAAAAT 524

FEATURE:
 US ID: 198 524 10991/

1 Sequence 10991 Application 35/1-198-524
 2 GENERAL INFORMATION:
 3 APPLICATION: Biosoft Inc
 4 TITLE OF INVENTION: New 1 Nucleic Acids and Polypeptides
 5 FILE REFERENCE: 21,274,614
 6 CURRENT APPLICATION NUMBER: 02/000,006
 7 PRIOR APPLICATION NUMBER: 02/000,006
 8 PRIOR FILING DATE: 2000-09-19
 9 PRIOR APPLICATION NUMBER: 02/000,006
 10 PRIOR FILING DATE: 2000-09-19
 11 NUMBER OF SEQ ID NOS: 1,246
 12 SOFTWARE: custom
 13 SEQ ID NO: 10991
 14 LENGTH: 1427
 15 TYPE: NA
 16 ORGANISM: Homo sapiens
 17 FEATURE:
 18 NAME/KEY: misc feature
 19 LOCATION: (1)-(1427)
 20 OTHER INFORMATION: n equals a, t, c, g, or r
 21 SEQ ID NO: 10991

Query Match
 Best Local Similarity 84.8% Score 17.89 Len 13 Length 1427%
 Matches 19 Conserved 0 Mismatches 2 Indels 0 Gaps 0

27 1 ATGAAAATGAAAATGAAAAT 21
 28 505 ATGAAAATGAAAATGAAAAT 500

FEATURE:
 US ID: 198 500 19776
 1 Sequence 1977 Application 35/1-198-500
 2 GENERAL INFORMATION:
 3 APPLICATION: Biosoft Inc
 4 TITLE OF INVENTION: Nucleic Acids, Proteins, and Antisense
 5 FILE REFERENCE: 21,274,614
 6 CURRENT APPLICATION NUMBER: 02/000,006
 7 PRIOR APPLICATION NUMBER: 02/000,006
 8 PRIOR FILING DATE: 2000-09-19
 9 PRIOR APPLICATION NUMBER: 02/000,006
 10 PRIOR FILING DATE: 2000-09-19
 11 NUMBER OF SEQ ID NOS: 1,246
 12 SOFTWARE: custom
 13 SEQ ID NO: 1977

[illegible]

100

[illegible][illegible]

Index

[illegible][illegible]

Figure 1

RESERVED
 CLASS 44—CLOTHING
 SWEATERS, T-SHIRTS, AND TURTLE NECK SHIRTS
 PATENT NO. 7,611,117
 GENERAL INFORMATION:
 APPLICANT: TOSHIBA, LIMITED
 APPLICANT'S ADDRESS: TOKYO, JAPAN
 ATTORNEY: LIPSCHUTZ, FRIEDMAN, LEVINE & LIPSCHUTZ, P.C.
 TITLE OF INVENTION: METHOD OF MANUFACTURING
 TURTLE NECK SHIRTS

[illegible]

COUNTRY: CANADA
 DATE: 04/11/2004
 ORDER REFERENCE: 1001
 ORDER NUMBER: 1001
 ORDER TYPE: REPLY ORDER
 ORDER DATE: 04/11/2004
 ORDER TIME: 10:24:41
 ORDERING SYSTEM: 1001/M2/00000000
 SUBORDINATE: 1001/M2/00000000
 CURRENT APPLICATION: 1001/M2/00000000
 APPLICATION NUMBER: 1001/M2/00000000

[illegible]

Library Method
Post Local Similarity Method
Matrices

1 APPLICANT: ROSS, PAUL
 2 APPLICANT: JARVIS, PAUL
 3 TITLE OF INVENTION: METHOD FOR TRANSDUCING ACOUSTIC MATERIALS
 4 TITLE OF INVENTION: METHOD FOR TRANSDUCING ACOUSTIC MATERIALS
 5 NUMBER OF SEQUENCES: 11
 6 CORRESPONDENCE ADDRESS:
 7 ADDRESS: NATIONAL RENEWABLE ENERGY LABORATORY
 8 STREET: 1417 GALE BUILDING
 9 CITY: Golden
 10 STATE: CO
 11 COUNTRY: U.S.A.
 12 ZIP: 80401-3995
 13 COMPUTER READABLE FORM:
 14 MULTIMEDIA TYPE: Pict 1.0
 15 MULTIMEDIA TYPE: Pict 1.0
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 100 MULTIMEDIA TYPE: Pict 1.0

101 100% Match
 102 Best Local Similarity: 90.7%
 103 Matches: 100, Unsuccessful: 0, Mismatches: 0, Indels: 0, Gaps: 0

104 100% Match
 105 Best Local Similarity: 90.7%
 106 Matches: 100, Unsuccessful: 0, Mismatches: 0, Indels: 0, Gaps: 0

107 100% Match
 108 Best Local Similarity: 90.7%
 109 Matches: 100, Unsuccessful: 0, Mismatches: 0, Indels: 0, Gaps: 0

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101 100% Match
 102 Best Local Similarity: 90.7%
 103 Matches: 100, Unsuccessful: 0, Mismatches: 0, Indels: 0, Gaps: 0

104 100% Match
 105 Best Local Similarity: 90.7%
 106 Matches: 100, Unsuccessful: 0, Mismatches: 0, Indels: 0, Gaps: 0

107 100% Match
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 109 Matches: 100, Unsuccessful: 0, Mismatches: 0, Indels: 0, Gaps: 0

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1. The first part of the document discusses the importance of maintaining accurate records of all transactions and activities. It emphasizes that proper record-keeping is essential for transparency and accountability, particularly in financial matters. The text outlines various methods for organizing and storing data, including digital databases and physical filing systems. It also mentions the need for regular audits and reviews to ensure the integrity of the information.

2. The second part of the document focuses on the role of communication in achieving organizational goals. It highlights the importance of clear and concise communication, both internally and externally. The text provides guidelines for effective communication, such as using appropriate language, listening actively, and providing feedback. It also discusses the benefits of open communication and how it can foster a collaborative work environment.

3. The third part of the document addresses the issue of time management. It recognizes that time is a valuable resource and that efficient use of time is crucial for productivity. The text offers several strategies for managing time, including prioritizing tasks, setting deadlines, and delegating responsibilities. It also emphasizes the importance of taking breaks and maintaining a healthy work-life balance to prevent burnout.

4. The fourth part of the document discusses the importance of continuous learning and development. It states that in a rapidly changing world, individuals and organizations must stay up-to-date with the latest trends and technologies. The text encourages a growth mindset and provides suggestions for acquiring new skills, such as attending workshops, taking courses, and seeking mentorship. It also mentions the importance of reflecting on one's own experiences and learning from mistakes.

5. The fifth part of the document concludes by summarizing the key points discussed throughout the document. It reiterates the importance of record-keeping, communication, time management, and continuous learning. The text expresses optimism about the future and encourages individuals and organizations to strive for excellence in all their endeavors.

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RESULT 2

US-09-057-570-2

1 Sequence 2, Application US/99/000000
2 Patent No. 6014266
3 GENERAL INFORMATION
4 APPLICANT: SCHOEN, ROBERT A.M.
5 TITLE OF INVENTION: Live attenuated Actinobacillus
6 TITLE OF INVENTION: pleuropneumoniae
7 NUMBER OF SEQUENCES: 7
8 ADDRESS: AKZO NO. 6014266 Patent Department
9 STREET: 1000 Riverside Drive, Suite 200
10 CITY: Rockville
11 STATE: Maryland
12 COUNTRY: USA
13 ZIP: 20850
14 MEDIUM TYPE: EMBRYO disk
15 COMPUTER: IBM PC compatible
16 OPERATING SYSTEM: MS-DOS 6.22
17 SOFTWARE: PATENT IN RELEASE #100, Version #1.30 (EP-9)
18 ALL INFORMATION SUBJECT TO: 6014266
19 FILING DATE: 09 APR 1998
20 PUBLICATION DATE: 09 APR 1998
21 ATTORNEY/AGENT INFORMATION:
22 NAME: Gentry, Mary E.
23 REGISTRATION NUMBER: 94459
24 TELEPHONE: (301) 948-9740
25 TELEFAX: (301) 948-9740
26 INFORMATION FOR SEQ. 1: Nucleotide
27 SEQUENCE CHARACTERISTICS:
28 LENGTH: 1657 amino acids
29 TYPE: amino acid
30 ORGANISM: Human
31 Molecule type: protein
32 US-09-057-570-2

Query Match: 6.444 Score 1100 Length 1657

Best Local Similarity: 21.984 Prod. No. 1111

Matched: 48 Conserved: 411 Mismatches: 114 Gaps: 17

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RESULT 4

US-09-057-570-7

1 Sequence 7, Application US/99/000000

2 Patent No. 6014266

3 GENERAL INFORMATION

4 APPLICANT: SCHOEN, ROBERT A.M.

5 TITLE OF INVENTION: Live attenuated Actinobacillus

6 TITLE OF INVENTION: pleuropneumoniae

7 NUMBER OF SEQUENCES: 7

8 ADDRESS: AKZO NO. 6014266 Patent Department

9 STREET: 1000 Riverside Drive, Suite 200

10 CITY: Rockville

11 STATE: Maryland

12 COUNTRY: USA

13 ZIP: 20850

14 MEDIUM TYPE: EMBRYO disk

15 COMPUTER: IBM PC compatible

16 OPERATING SYSTEM: MS-DOS 6.22

17 SOFTWARE: PATENT IN RELEASE #100, Version #1.30 (EP-9)

18 ALL INFORMATION SUBJECT TO: 6014266

19 FILING DATE: 09 APR 1998

20 PUBLICATION DATE: 09 APR 1998

21 ATTORNEY/AGENT INFORMATION:

22 NAME: Gentry, Mary E.

23 REGISTRATION NUMBER: 94459

24 TELEPHONE: (301) 948-9740

25 TELEFAX: (301) 948-9740

26 INFORMATION FOR SEQ. 1: Nucleotide

27 SEQUENCE CHARACTERISTICS:

28 LENGTH: 1805 amino acids

29 TYPE: amino acid

30 ORGANISM: Human

31 Molecule type: protein

US-09-057-570-7

Query Match: 6.444 Score 1100 Length 1657

Best Local Similarity: 21.984 Prod. No. 1111

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Sequence 13, Affixation: 007/04020000

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Sequence 17, Affixation: 007/04020000

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Sequence 22, Affixation: 007/04020000

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Sequence 24, Affixation: 007/04020000

Sequence 25, Affixation: 007/04020000

Sequence 26, Affixation: 007/04020000

Sequence 27, Affixation: 007/04020000

Sequence 28, Affixation: 007/04020000

Sequence 29, Affixation: 007/04020000

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Sequence 55, Affixation: 007/04020000

Sequence 56, Affixation: 007/04020000

Sequence 57, Affixation: 007/04020000

Sequence 58, Affixation: 007/04020000


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1  APPLICANT: Knechtley, Phillip
2  APPLICANT: Knechtley, Phillip
3  TITLE OF INVENTION: GENETIC DATA SEQUENCES OF ASHURA (POSSYPH)
4  TITLE OF INVENTION: AND USES THEREOF
5  NUMBER OF SEQUENCES: 1752
6  PRESIDENTIAL ADDRESS:
7  APPLICANT: Knechtley, Phillip
8  STREET: 4054 Cornwellis Road
9  CITY: Research Triangle Park
10 STATE: NO. 620264th Carolina
11 COUNTRY: USA
12 ZIP: 27709
13 NUMBER BLANK PAGE:
14 MEDIUM TYPE: floppy disk
15 COMPUTER: IBM pc compatible
16 SOFTWARE: Patent in Release #1.0, Version #1.30
17 CURRENT APPLICATION DATA:
18 ADDITIONAL NUMBER: 02/06/99, 416
19 FILING DATE: 24 Dec 1997
20 CLASSIFICATION: 4.0
21 PRIOR APPLICATION DATA:
22 APPLICATION NUMBER: 08-09-094
23 FILING DATE: 31 Dec 1996
24 ATTORNEY/AGENT INFORMATION:
25 NAME: Mohr, J. Timothy
26 REGISTRATION NUMBER: 06,241
27 REFERENCE TO PRIOR APPLICATIONS:
28 TELEPHONE: 919 541 8587
29 TELEFAX: 919 541 8587
30 E-MAIL FOR SEQ ID NO: 288:
31 SEQUENCE CHARACTERISTICS:
32 LENGTH: 607 base pairs
33 TYPE: nucleic acid
34 SUBSTANCE: single
35 MOLECULE TYPE: DNA (genomic)
36 ORGANISM: PASTORAL
37 US 08 098 416 186

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Query Match: 4.76% Score 61.96 E-11.00 P-1.00

Best Local Similarity: 47.86% Prod. No. 1.00

Mismatch: 214 Conservative 0 Mismatched 186 Indels 10 Gaps 6

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37 106 TAAATATATATATATATATATATATATATATATATATATATATATAT 317
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39 106 TAAATATATATATATATATATATATATATATATATATATATATATAT 349
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RESULT 6
US 08 998 416 186
: Sequence ID: Application US/08 09416
: Patent No. 620264
: GENERAL INFORMATION:
: APPLICANT: Knechtley, Phillip
: APPLICANT: Knechtley, Phillip
: APPLICANT: Knechtley, Phillip
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: NUMBER OF SEQUENCES: 1752
: PRESIDENTIAL ADDRESS:
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: E-MAIL FOR SEQ ID NO: 186:
: SEQUENCE CHARACTERISTICS:
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: SUBSTANCE: single
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: ORGANISM: PASTORAL
: US 08 998 416 186

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45 106 TAAATATATATATATATATATATATATATATATATATATATATATAT 445
46 111 111 111 111 111 111 111 111 111 111 111 111 111 111

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Figure 1 consists of 12 panels (a-l) showing Western blot analysis of protein expression in 293T cells. The panels are arranged in two columns. The left column (a, c, e, g, i, k, m) shows the expression of various proteins, and the right column (b, d, f, h, j, l, n) shows the expression of the corresponding siRNA target proteins. The proteins are: p53 (a, b), p21 (c, d), p16 (e, f), p14 (g, h), p15 (i, j), p18 (k, l), p19 (m, n), p20 (o, p), p21 (q, r), p22 (s, t), p23 (u, v), and p24 (w, x). Molecular weight markers are indicated on the right of each panel.

[illegible]

Figure	Figure	Figure
Fig. 1	Fig. 2	Fig. 3
Fig. 4	Fig. 5	Fig. 6
Fig. 7	Fig. 8	Fig. 9
Fig. 10	Fig. 11	Fig. 12
Fig. 13	Fig. 14	Fig. 15
Fig. 16	Fig. 17	Fig. 18
Fig. 19	Fig. 20	Fig. 21
Fig. 22	Fig. 23	Fig. 24
Fig. 25	Fig. 26	Fig. 27
Fig. 28	Fig. 29	Fig. 30
Fig. 31	Fig. 32	Fig. 33
Fig. 34	Fig. 35	Fig. 36
Fig. 37	Fig. 38	Fig. 39
Fig. 40	Fig. 41	Fig. 42
Fig. 43	Fig. 44	Fig. 45
Fig. 46	Fig. 47	Fig. 48
Fig. 49	Fig. 50	Fig. 51
Fig. 52	Fig. 53	Fig. 54
Fig. 55	Fig. 56	Fig. 57
Fig. 58	Fig. 59	Fig. 60
Fig. 61	Fig. 62	Fig. 63
Fig. 64	Fig. 65	Fig. 66
Fig. 67	Fig. 68	Fig. 69
Fig. 70	Fig. 71	Fig. 72
Fig. 73	Fig. 74	Fig. 75
Fig. 76	Fig. 77	Fig. 78
Fig. 79	Fig. 80	Fig. 81
Fig. 82	Fig. 83	Fig. 84
Fig. 85	Fig. 86	Fig. 87
Fig. 88	Fig. 89	Fig. 90
Fig. 91	Fig. 92	Fig. 93
Fig. 94	Fig. 95	Fig. 96
Fig. 97	Fig. 98	Fig. 99
Fig. 100	Fig. 101	Fig. 102
Fig. 103	Fig. 104	Fig. 105
Fig. 106	Fig. 107	Fig. 108
Fig. 109	Fig. 110	Fig. 111
Fig. 112	Fig. 113	Fig. 114
Fig. 115	Fig. 116	Fig. 117
Fig. 118	Fig. 119	Fig. 120
Fig. 121	Fig. 122	Fig. 123
Fig. 124	Fig. 125	Fig. 126
Fig. 127	Fig. 128	Fig. 129
Fig. 130	Fig. 131	Fig. 132
Fig. 133	Fig. 134	Fig. 135
Fig. 136	Fig. 137	Fig. 138
Fig. 139	Fig. 140	Fig. 141
Fig. 142	Fig. 143	Fig. 144
Fig. 145	Fig. 146	Fig. 147
Fig. 148	Fig. 149	Fig. 150
Fig. 151	Fig. 152	Fig. 153
Fig. 154	Fig. 155	Fig. 156
Fig. 157	Fig. 158	Fig. 159
Fig. 160	Fig. 161	Fig. 162
Fig. 163	Fig. 164	Fig. 165
Fig. 166	Fig. 167	Fig. 168
Fig. 169	Fig. 170	Fig. 171
Fig. 172	Fig. 173	Fig. 174
Fig. 175	Fig. 176	Fig. 177
Fig. 178	Fig. 179	Fig. 180
Fig. 181	Fig. 182	Fig. 183
Fig. 184	Fig. 185	Fig. 186
Fig. 187	Fig. 188	Fig. 189
Fig. 190	Fig. 191	Fig. 192
Fig. 193	Fig. 194	Fig. 195
Fig. 196	Fig. 197	Fig. 198
Fig. 199	Fig. 200	Fig. 201
Fig. 202	Fig. 203	Fig. 204
Fig. 205	Fig. 206	Fig. 207
Fig. 208	Fig. 209	Fig. 210
Fig. 211	Fig. 212	Fig. 213
Fig. 214	Fig. 215	Fig. 216
Fig. 217	Fig. 218	Fig. 219
Fig. 220	Fig. 221	Fig. 222
Fig. 223	Fig. 224	Fig. 225
Fig. 226	Fig. 227	Fig. 228
Fig. 229	Fig. 230	Fig. 231
Fig. 232	Fig. 233	Fig. 234
Fig. 235	Fig. 236	Fig. 237
Fig. 238	Fig. 239	Fig. 240
Fig. 241	Fig. 242	Fig. 243
Fig. 244	Fig. 245	Fig. 246
Fig. 247	Fig. 248	Fig. 249
Fig. 250	Fig. 251	Fig. 252
Fig. 253	Fig. 254	Fig. 255
Fig. 256	Fig. 257	Fig. 258
Fig. 259	Fig. 260	Fig. 261
Fig. 262	Fig. 263	Fig. 264
Fig. 265	Fig. 266	Fig. 267
Fig. 268	Fig. 269	Fig. 270
Fig. 271	Fig. 272	Fig. 273
Fig. 274	Fig. 275	Fig. 276
Fig. 277	Fig. 278	Fig. 279
Fig. 280	Fig. 281	Fig. 282
Fig. 283	Fig. 284	Fig. 285
Fig. 286	Fig. 287	Fig. 288
Fig. 289	Fig. 290	Fig.

[illegible]

Figure 1: Schematic representation of the experimental design. The figure shows a flowchart of the experimental design. It starts with 'Pretest' leading to 'Main Experiment'. The 'Main Experiment' is divided into 'Condition 1' and 'Condition 2'. 'Condition 1' leads to 'Posttest' and 'Debriefing'. 'Condition 2' leads to 'Posttest' and 'Debriefing'. The 'Posttest' is a 'Posttest' and the 'Debriefing' is a 'Debriefing'.

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	(a)	(b)	(c)	(d)	(e)	(f)	(g)	(h)	(i)	(j)	(k)	(l)	(m)	(n)	(o)	(p)	(q)	(r)	(s)	(t)	(u)	(v)	(w)	(x)	(y)	(z)	(aa)	(ab)	(ac)	(ad)	(ae)	(af)	(ag)	(ah)	(ai)	(aj)	(ak)	(al)	(am)	(an)	(ao)	(ap)	(aq)	(ar)	(as)	(at)	(au)	(av)	(aw)	(ax)	(ay)	(az)	(ba)	(bb)	(bc)	(bd)	(be)	(bf)	(bg)	(bh)	(bi)	(bj)	(bk)	(bl)	(bm)	(bn)	(bo)	(bp)	(bq)	(br)	(bs)	(bt)	(bu)	(bv)	(bw)	(bx)	(by)	(bz)	(ca)	(cb)	(cc)	(cd)	(ce)	(cf)	(cg)	(ch)	(ci)	(cj)	(ck)	(cl)	(cm)	(cn)	(co)	(cp)	(cq)	(cr)	(cs)	(ct)	(cu)	(cv)	(cw)	(cx)	(cy)	(cz)	(da)	(db)	(dc)	(dd)	(de)	(df)	(dg)	(dh)	(di)	(dj)	(dk)	(dl)	(dm)	(dn)	(do)	(dp)	(dq)	(dr)	(ds)	(dt)	(du)	(dv)	(dw)	(dx)	(dy)	(dz)	(ea)	(eb)	(ec)	(ed)	(ee)	(ef)	(eg)	(eh)	(ei)	(ej)	(ek)	(el)	(em)	(en)	(eo)	(ep)	(eq)	(er)	(es)	(et)	(eu)	(ev)	(ew)	(ex)	(ey)	(ez)	(fa)	(fb)	(fc)	(fd)	(fe)	(ff)	(fg)	(fh)	(fi)	(fj)	(fk)	(fl)	(fm)	(fn)	(fo)	(fp)	(fq)	(fr)	(fs)	(ft)	(fu)	(fv)	(fw)	(fx)	(fy)	(fz)	(ga)	(gb)	(gc)	(gd)	(ge)	(gf)	(gg)	(gh)	(gi)	(gj)	(gk)	(gl)	(gm)	(gn)	(go)	(gp)	(gq)	(gr)	(gs)	(gt)	(gu)	(gv)	(gw)	(gx)	(gy)	(gz)	(ha)	(hb)	(hc)	(hd)	(he)	(hf)	(hg)	(hh)	(hi)	(hj)	(hk)	(hl)	(hm)	(hn)	(ho)	(hp)	(hq)	(hr)	(hs)	(ht)	(hu)	(hv)	(hw)	(hx)	(hy)	(hz)	(ia)	(ib)	(ic)	(id)	(ie)	(if)	(ig)	(ih)	(ii)	(ij)	(ik)	(il)	(im)	(in)	(io)	(ip)	(iq)	(ir)	(is)	(it)	(iu)	(iv)	(iw)	(ix)	(iy)	(iz)	(ja)	(jb)	(jc)	(jd)	(je)	(jf)	(jg)	(jh)	(ji)	(jj)	(jk)	(jl)	(jm)	(jn)	(jo)	(jp)	(jq)	(jr)	(js)	(jt)	(ju)	(jv)	(jw)	(jx)	(jy)	(jz)	(ka)	(kb)	(kc)	(kd)	(ke)	(kf)	(kg)	(kh)	(ki)	(kj)	(kk)	(kl)	(km)	(kn)	(ko)	(kp)	(kq)	(kr)	(ks)	(kt)	(ku)	(kv)	(kw)	(kx)	(ky)	(kz)	(la)	(lb)	(lc)	(ld)	(le)	(lf)	(lg)	(lh)	(li)	(lj)	(lk)	(ll)	(lm)	(ln)	(lo)	(lp)	(lq)	(lr)	(ls)	(lt)	(lu)	(lv)	(lw)	(lx)	(ly)	(lz)	(ma)	(mb)	(mc)	(md)	(me)	(mf)	(mg)	(mh)	(mi)	(mj)	(mk)	(ml)	(mm)	(mn)	(mo)	(mp)	(mq)	(mr)	(ms)	(mt)	(mu)	(mv)	(mw)	(mx)	(my)	(mz)	(na)	(nb)	(nc)	(nd)	(ne)	(nf)	(ng)	(nh)	(ni)	(nj)	(nk)	(nl)	(nm)	(nn)	(no)	(np)	(nq)	(nr)	(ns)	(nt)	(nu)	(nv)	(nw)	(nx)	(ny)	(nz)	(oa)	(ob)	(oc)	(od)	(oe)	(of)	(og)	(oh)	(oi)	(oj)	(ok)	(ol)	(om)	(on)	(oo)	(op)	(oq)	(or)	(os)	(ot)	(ou)	(ov)	(ow)	(ox)	(oy)	(oz)	(pa)	(pb)	(pc)	(pd)	(pe)	(pf)	(pg)	(ph)	(pi)	(pj)	(pk)	(pl)	(pm)	(pn)	(po)	(pp)	(pq)	(pr)	(ps)	(pt)	(pu)	(pv)	(pw)	(px)	(py)	(pz)	(qa)	(qb)	(qc)	(qd)	(qe)	(qf)	(qg)	(qh)	(qi)	(qj)	(qk)	(ql)	(qm)	(qn)	(qo)	(qp)	(qq)	(qr)	(qs)	(qt)	(qu)	(qv)	(qw)	(qx)	(qy)	(qz)	(ra)	(rb)	(rc)	(rd)	(re)	(rf)	(rg)	(rh)	(ri)	(rj)	(rk)	(rl)	(rm)	(rn)	(ro)	(rp)	(rq)	(rr)	(rs)	(rt)	(ru)	(rv)	(rw)	(rx)	(ry)	(rz)	(sa)	(sb)	(sc)	(sd)	(se)	(sf)	(sg)	(sh)	(si)	(sj)	(sk)	(sl)	(sm)	(sn)	(so)	(sp)	(sq)	(sr)	(ss)	(st)	(su)	(sv)	(sw)	(sx)	(sy)	(sz)	(ta)	(tb)	(tc)	(td)	(te)	(tf)	(tg)	(th)	(ti)	(tj)	(tk)	(tl)	(tm)	(tn)	(to)	(tp)	(tq)	(tr)	(ts)	(tt)	(tu)	(tv)	(tw)	(tx)	(ty)	(tz)	(ua)	(ub)	(uc)	(ud)	(ue)	(uf)	(ug)	(uh)	(ui)	(uj)	(uk)	(ul)	(um)	(un)	(uo)	(up)	(uq)	(ur)	(us)	(ut)	(uu)	(uv)	(uw)	(ux)	(uy)	(uz)	(va)	(vb)	(vc)	(vd)	(ve)	(vf)	(vg)	(vh)	(vi)	(vj)	(vk)	(vl)	(vm)	(vn)	(vo)	(vp)	(vq)	(vr)	(vs)	(vt)	(vu)	(vv)	(vw)	(vx)	(vy)	(vz)	(wa)	(wb)	(wc)	(wd)	(we)	(wf)	(wg)	(wh)	(wi)	(wj)	(wk)	(wl)	(wm)	(wn)	(wo)	(wp)	(wq)	(wr)	(ws)	(wt)	(wu)	(wv)	(ww)	(wx)	(wy)	(wz)	(xa)	(xb)	(xc)	(xd)	(xe)	(xf)	(xg)	(xh)	(xi)	(xj)	(xk)	(xl)	(xm)	(xn)	(xo)	(xp)	(xq)	(xr)	(xs)	(xt)	(xu)	(xv)	(xw)	(xx)	(xy)	(xz)	(ya)	(yb)	(yc)	(yd)	(ye)	(yf)	(yg)	(yh)	(yi)	(yj)	(yk)	(yl)	(ym)	(yn)	(yo)	(yp)	(yq)	(yr)	(ys)	(yt)	(yu)	(yv)	(yw)
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1	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
2	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
3	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
4	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
5	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81																			

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Pos. 1	No.	Series	Ref.	Length	Th	Sec	Lat	Long	Alt
1	1018	99.7	7.4	7.4	7.4	7.4	7.4	7.4	7.4
2	1006.6	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
3	87.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
4	86.38	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
5	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
6	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
7	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
8	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
9	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
10	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
11	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
12	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
13	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
14	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
15	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
16	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
17	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
18	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
19	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
20	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
21	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
22	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
23	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
24	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
25	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
26	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
27	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
28	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
29	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4
30	86.2	4.1	7.4	7.4	7.4	7.4	7.4	7.4	7.4



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